=> FILE REGISTRY

=> Uploading C:\Program Files\STNEXP\Queries\10585478_01252010_2.str

1 2 3 4 5 chain bonds: 1-2 2-3 3-4 3-5 exact/norm bonds: 1-2 exact bonds: 2-3 3-4 3-5

Match level: 1:Atom 2:CLASS 3:CLASS 4:CLASS 5:CLASS Generic attributes: 1: Number of Hetero Atoms: 2 or more

L1 STRUCTURE UPLOADED

=> d L1 HAS NO ANSWERS L1 STR

Structure attributes must be viewed using STN Express query preparation.

=> s 11 full FULL SEARCH INITIATED 06:46:21 FILE 'REGISTRY' FULL SCREEN SEARCH COMPLETED - 19112 TO ITERATE

100.0% PROCESSED 19112 ITERATIONS SEARCH TIME: 00.00.01 1000 ANSWERS

L3

Uploading C:\Program Files\STNEXP\Queries\10585478 01252010 4.str -[-]1-6.

chain nodes : 1 2 3 4 5 chain bonds : 1-2 2-3 3-4 3-5 exact/norm bonds : 1 - 2exact bonds :

2-3 3-4 3-5

Match level : 1:Atom 2:CLASS 3:CLASS 4:CLASS 5:CLASS Generic attributes : Saturation : Unsaturated Number of Hetero Atoms : 2 or more

L8 STRUCTURE UPLOADED

=> d L8 HAS NO ANSWERS L8 STR



Structure attributes must be viewed using STN Express query preparation.

=> s 18 sss sub=16 full FULL SUBSET SEARCH INITIATED 06:49:34 FILE 'REGISTRY' FULL SUBSET SCREEN SEARCH COMPLETED -520 TO ITERATE

100.0% PROCESSED 488 ANSWERS 520 ITERATIONS SEARCH TIME: 00.00.01

488 SEA SUB=L6 SSS FUL L8

=> s 110 and caplus/lc 69979096 CAPLUS/LC

287 L10 AND CAPLUS/LC

=> Uploading C:\Program Files\STNEXP\Queries\10585478_01252010_5.str



Match level:
1:Atom 2:CLASS 3:CLASS 4:CLASS 5:CLASS 6:CLASS
Generic attributes:
1:

Saturation : Unsaturated Number of Hetero Atoms : 2 or more

L13 STRUCTURE UPLOADED

=> d L13 HAS NO ANSWERS L13 STR Hy

=> s 113 sss sub=110 full FULL SUBSET SEARCH INITIATED 06:50:47 FILE 'REGISTRY' FULL SUBSET SCREEN SEARCH COMPLETED - 153 TO ITERATE

100.0% PROCESSED 153 ITERATIONS SEARCH TIME: 00.00.01 132 ANSWERS

L15 132 SEA SUB=L10 SSS FUL L13

=> s 115 and caplus/1c

```
69979096 CAPLUS/LC
132 L15 AND CAPLUS/LC
```

=> file caplus

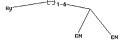
=> s 116 L17 10 L16

=> d 117 ibib qi abs hitstr 1-10

=>

1.16

Uploading C:\Program Files\STNEXP\Queries\10585478_01252010_6.str



3

chain nodes:
1 2 3 4 5
chain bonds:
1-2 2-3 3-4 3-5
exact/norm bonds:
1-2
exact bonds:
2-3 3-4 3-5

Saturation

Match level: 1:Atom 2:CLASS 3:CLASS 4:CLASS 5:CLASS Generic attributes:

Number of Hetero Atoms : 2 or more

: Unsaturated

L19 STRUCTURE UPLOADED

=> d L19 HAS NO ANSWERS L19 STR

ĆN

=> s 119 sss sub=118 full FULL SUBSET SEARCH INITIATED 07:00:43 FILE 'REGISTRY' FULL SUBSET SCREEN SEARCH COMPLETED - 742 TO ITERATE

=> logoff hold

L21 690 SEA SUB=L18 SSS FUL L19

```
Uploading C:\Program Files\STNEXP\Queries\10585478_01252010_3.str
                        CN
                                 CN.
chain nodes :
1 2 3 4
ring nodes :
7 8 9 10 11
chain bonds :
1-2 1-7 2-3 2-4
ring bonds :
7-8 7-11 8-9 9-10 10-11
exact/norm bonds :
1-7 7-8 7-11 8-9 9-10 10-11
exact bonds :
1-2 2-3 2-4
Match level :
1:CLASS 2:CLASS 3:CLASS 4:CLASS 7:CLASS 8:Atom 9:Atom 10:Atom 11:Atom
L22
      STRUCTURE UPLOADED
=> s 122 sss sub=121 full
FULL SUBSET SEARCH INITIATED 07:01:31 FILE 'REGISTRY'
FULL SUBSET SCREEN SEARCH COMPLETED -
                                         690 TO ITERATE
100.0% PROCESSED
                     690 ITERATIONS
                                                             287 ANSWERS
SEARCH TIME: 00.00.01
           287 SEA SUB=L21 SSS FUL L22
=> file caplus
=> s 124/USES
            83 L24
       7936248 USES/RL
L25
           13 L24/USES
                (L24 (L) USES/RL)
=> d 125 ibib qi abs hitstr 1-13
```